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## **Inheritance of ectopic ureters in Entlebucher Mountain Dogs**

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**Abstract:** To test the hypothesis of a heritable base of ectopic ureters (EU) in Entlebucher Mountain Dogs (EMD) and to elucidate associated risk factors and mode of inheritance of the disease, 565 EMD were clinically investigated and population genetic analyses performed. Based on the location of the most caudal termination of the ureteral openings, 552 EMD were classified into three phenotype groups trigone, intravesically and extravesically ectopic based on results of abdominal sonography, urethra-cystoscopy and/or contrast-enhanced computed tomography. One-third (32.9%) of the phenotyped animals had normal terminations of both ureters in the bladder trigone, 47.3% had at least one intravesicular ectopic termination and 19.8% had at least one extravesicular ectopic termination. Multivariate mixed logistic regression revealed gender as a risk factor associated with EU as males were more often affected than females. Complex segregation analysis indicated a hereditary basis for EU in EMD and the involvement of a major gene in the occurrence of the extravesicular EU phenotype.

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12 Keywords  
13 Ureteral ectopia, population genetics, canine, breeding selection, ureteral orifice

## 14 Summary

15 To test the hypothesis of a heritable base of ectopic ureters (EU) in Entlebucher Mountain  
16 Dogs (EMD) and to elucidate associated risk factors and mode of inheritance of the disease,  
17 565 EMD were clinically investigated and population genetic analyses performed. Based on  
18 the location of the most caudal termination of the ureteral openings, 552 EMD were classified  
19 into three phenotype groups trigone, intravesically and extravesically ectopic based on results  
20 of abdominal sonography, urethra-cystoscopy and/or contrast-enhanced computed  
21 tomography. One third (32.9%) of the phenotyped animals had normal terminations of both  
22 ureters in the bladder trigone, 47.3% had at least one intravesicular ectopic termination and  
23 19.8% had at least one extravesicular ectopic termination. Mixed multivariable logistic  
24 regression revealed gender as a risk factor associated with EU as males were more often  
25 affected than females. Complex segregation analysis indicated a hereditary basis for EU in  
26 EMD and the involvement of a major gene in the occurrence of the extravesicular EU  
27 phenotype.

## 28 Introduction

29 Ureteral ectopia is a congenital abnormality in which one or both ureters terminate in a  
30 position other than the trigone of the urinary bladder (Osborne *et al.* 1995). Ectopic ureters  
31 (EU) are classified according to the location of their termination as intravesicular (IVEU) or  
32 extravesicular (EVEU) (North *et al.* 2010), and according to their course until their terminal  
33 orifice as intramural or extramural (Ho *et al.* 2011).

34 In the dog, the condition is generally rare with reported incidences below 0.05% for clinically  
35 apparent cases (Hayes 1974; Smith *et al.* 1981; Dean *et al.* 1988). Entlebucher Mountain  
36 Dogs (EMD), Briards, Bulldogs, Golden Retrievers, Labrador Retrievers, Griffons, Border  
37 Terriers, Fox Terriers, Skye Terriers, West Highland White Terriers, Siberian Huskies,  
38 Newfoundland dogs, Miniature and Toy Poodles have been found to be at an increased risk  
39 for EU (Hayes 1974; Hayes 1984; Holt *et al.* 2000; Eckrich Specker 2006; North *et al.* 2010;

40 Bitterli 2011). The most common clinical sign is the occurrence of urinary incontinence at a  
41 young age. However, some animals, and especially males, do not show clinical symptoms  
42 until they reach an advanced age (Holt 1990, Holt & Moore 1995). In a case series of 50 dogs  
43 undergoing EU surgery (Reichler et al., 2012) urinary incontinence was observed for the first  
44 time within the first year of life in 90% of the females but only in 50% of the males. 16% of  
45 the males were older than 5 years when the owner observed urinary incontinence for the first  
46 time (unpublished data). Besides incontinence, recurrent urinary tract infections (e.g. cystitis,  
47 pyelonephritis), hydroureter and hydronephrosis are of clinical importance (Stone & Mason  
48 1990; Cannizzo *et al.* 2003). In severely affected dogs, the course of the disease may be fatal.  
49 The hypothesis of a genetic background of EU is supported by reports of familial  
50 aggregations (Johnston *et al.* 1977; Holt *et al.* 1982). Familial increased incidence of EU has  
51 also been reported in humans (Deweerd & Feeney 1967; Musselman & Barry 1973). Reports  
52 in related EMD and an overrepresentation of the breed in surgical EU cases (Eckrich Specker  
53 2006; North *et al.* 2010; Bitterli 2011; Reichler *et al.* 2012) led to the implementation of a  
54 screening program for EU with the participation of the Swiss, German, Dutch and Austrian  
55 EMD kennel clubs (SKES, SSV-ES, ESC and VSSÖ). Starting in mid-2008, the German  
56 kennel club required breeding dogs to be examined for presence of EU, whereas the Swiss,  
57 Dutch and Austrian kennel club strongly recommended testing. A restrictive program which  
58 excluded severely affected dogs was established in Germany in 2009. However, the high  
59 incidence of EU in the examined EMD (n= 308, 45% IVEU and 19% EVEU) (Bitterli 2011)  
60 together with an overall small population size, the effective population size in Switzerland  
61 being 32 in 2009 (Staub 2011), make the establishment of a reasonable breeding strategy  
62 rather difficult. The knowledge of the mode of inheritance improves the chance to identify  
63 causative genes (Snow & Wijnsman 1998).

64 The objective of this study was to evaluate risk factors associated with the occurrence of EU  
65 in EMD and to assess its mode of inheritance and its heritability.

## 66 Materials and Methods

67 Data: This multi-center, cross-sectional study of kennel club registered EMD was approved  
68 by the Swiss Federal Veterinary Office. Six institutes with board certified radiologists or  
69 internists in Switzerland, Germany, Austria and the Netherlands participated in the study.  
70 Pedigree information was provided by the kennels and every dog was identified by its  
71 microchip. An intravenous catheter was placed in the cephalic vein and a blood sample was  
72 collected into EDTA tubes and stored at  $-20^{\circ}\text{C}$  for future molecular studies. Abdominal  
73 sonography with or without sedation was the standard screening procedure. Visible jets of  
74 urine through the ureteral openings were documented separately in longitudinal and  
75 transverse planes in B-Mode (Lamb & Gregory 1998), Color Doppler or Color B-flow mode.  
76 Intravenous (IV) crystalloid infusion (lactated Ringer's solution) at a rate of 10ml/kg body  
77 weight (BW) and furosemide at 1mg/kg BW IV were added to reinforce the jet phenomenon.  
78 The proximal boundary of the urethra was defined as the point distal to the bladder, from  
79 where the diameter of the urinary tract remained constant (Rozear & Tidwell 2003). A normal  
80 distance between the ureteral openings and the vesicourethral junction was expected to be at  
81 least 1.5 to 2.0 cm dependent on the size of the dog. Smaller distances were considered to be  
82 ureteral orifice terminations in the "bladder neck". If the ureteral openings could not be  
83 convincingly localized and/or separately recorded, sonography was followed by contrast-  
84 enhanced computed tomography (CT excretory urography) or urethro-cystoscopy given the  
85 owner's consent for general anesthesia and examination. All exam-reports from the  
86 participating institutes were reviewed by the project group in Zurich and classified according  
87 to the following system: Normal, if the ureteral orifice terminated at the "bladder trigone",  
88 ectopic intravesicular, if it terminated at the bladder neck and ectopic extravesicular, if it  
89 terminated at the urethra which also included openings just at the vesicourethral junction.  
90 According to the more caudal location of the right or left ureteral orifice, each dog was

91 assigned one of four phenotypes: phenotype trigone, phenotype IVEU, phenotype EVEU or  
 92 unknown phenotype.

93 Statistical analysis: Descriptive statistics were calculated for the variables age at diagnosis,  
 94 gender, reproduction status, year of birth, season of birth, litter size, sex ratio and early-death.  
 95 Information on the last three variables was only available for Swiss and German EMD.

96 Season 1 was defined as birth between 1<sup>st</sup> December and the last day of February, Season 2  
 97 between 1<sup>st</sup> March and 31<sup>st</sup> May, Season 3 between 1<sup>st</sup> June and 31<sup>st</sup> August and Season 4  
 98 between 1<sup>st</sup> September and 30<sup>th</sup> November. Litter size was defined as the total number of  
 99 puppies born in a litter including stillborn pups. Sex ratio was calculated as the number of  
 100 male puppies divided by the total number of puppies per litter. The variable early-death  
 101 includes stillborn littermates or puppies that died during their first 8 weeks of life.

102 Associations of EU with categorical variables were analyzed using contingency tables.

103 Continuous variables were first checked for normality using QQ plots. If a variable was not  
 104 normally distributed an appropriate non-parametric test was performed. Gender distribution of  
 105 early-death puppies within the litters was evaluated against a ratio of 1:1. Additionally, an  
 106 expanded dataset of the last 12 years (2000-2011) for gender of all born and all early-death  
 107 puppies was examined. In order to evaluate the consequences of breeding restriction, EMD  
 108 born before the start of the screening program (2006 - 2007) were compared to those born  
 109 after breeding restrictions were established (2009 -2010).

110 An univariate analysis with the categorical variables gender, early death, season, birth after  
 111 2008 and reproduction status; and the continuous variables litter size, sex ratio, year of birth  
 112 and age at diagnosis was performed on the dataset of Swiss and German EMD. A mixed  
 113 logistic regression model was examined to take correlated outcomes of littermates into  
 114 account. Three versions with binary encoding of the phenotypes were run: (i) trigone as  
 115 unaffected, IVEU together with EVEU as affected; (ii) trigone together with IVEU as  
 116 unaffected, EVEU as affected; and (iii) trigone as unaffected and EVEU as affected while

117 specifying IVEU as no information. A multivariate mixed logistic regression was performed  
 118 to determine risk factors associated with the three phenotypes. Potential interactions were also  
 119 evaluated. Each variable was added separately to the model and its effect evaluated on the  
 120 basis of AIC (Akaike 1974). Each of the fixed effect variables in the multivariate model was  
 121 then evaluated based on its computed  $P$ -value. Only variables with significant effects were  
 122 kept in the final model. Log odds were used to measure the association between each  
 123 categorical variable and phenotypes. Descriptive statistics were performed with standard  
 124 software (IBM SPSS Statistics®, version 19.0 for Mac, SPSS Inc, Chicago, IL, USA),  
 125 whereas for the logistic regression models another software package was used (R: A language  
 126 and environment for statistical computing, R Foundation for Statistical Computing, Vienna,  
 127 Austria). Values of  $P < 0.05$  were considered significant.

128 Segregation analysis: Complex segregation analysis of the EU trait was carried out using PAP  
 129 (Pedigree Analysis Package 5.ed., Hasstedt 2002). EU was encoded as a dichotomous trait  
 130 with different combinations of the three phenotypes in three datasets (Table 3). The  
 131 prevalence of phenotypes was based on our screening results and calculated for the different  
 132 phenotype groupings within the datasets. Maximum likelihood procedures were used to  
 133 estimate the following parameters: allele frequency, transmission probabilities, dominance,  
 134 displacement and heritability. Five different models (general, environmental, mixed, major  
 135 gene and polygenic inheritance) were compared according to their hierarchy based on the  
 136 difference between their likelihoods ( $-2 \ln L$ ), the distribution of which follows a  $\chi^2$   
 137 distribution with degrees of freedom equal to the difference in the number of the parameters  
 138 estimated. A difference between two models was considered significant if  $P \leq 0.05$ .

139 Heritability: To estimate heritability of the EVEU trait the program MTDFREML (Boldman  
 140 *et al.* 1995) was used on an expanded dataset with all EMD that had a trigone or EVEU  
 141 phenotype in 2012, their ancestors of five generations and individuals to connect families.

142 Results

143 Out of 565 EMD (288 females, 277 males) one third (32 %) had normal terminations of both  
 144 ureters in the bladder trigone area, 46 % had at least one IVEU and 19 % had at least one  
 145 EVEU termination, whereas 13 individuals could not be classified (Table 1). There was a  
 146 significant difference in gender distribution with more males having EVEU than females ( $P <$   
 147  $0.001$ ). No difference in gender distribution was found within the early-death puppies ( $n = 63$ ,  
 148  $P = 0.450$ ) and the same was true when considering all born or all early death puppies within  
 149 the last 12 years ( $n = 3820$ ,  $P = 0.116$  vs.  $n = 274$ ,  $P = 0.589$ ). There was a significant change  
 150 in the distribution of phenotypes when comparing dogs born in the two years (2006 – 2007)  
 151 before the screening started in middle of 2008 with dogs born in the first two years after  
 152 breeding restrictions had been established (2009 – 2010;  $P < 0.001$ ; Figure 1). The percentage  
 153 of affected dogs changed from 49% and 25% to 50% and 10% for IVEU and EVEU,  
 154 respectively.

155 In all three versions (i, ii, iii) of the mixed multivariate logistic regression models  
 156 with the phenotyped Swiss and German EMD ( $n = 430$ ) gender was a supporting covariate  
 157 according to the best-fit models and the female gender was found to decrease the log odds  
 158 ratio. Additionally in version (ii) the covariates year of birth, season and age at diagnosis, and  
 159 in version (iii) birth after 2008 and season 3 were significantly associated with the outcome  
 160 (Table 2). No improvement was achieved when adding interactions or litters as a random  
 161 factor.

162 For the segregation analysis of the three datasets 282 phenotyped (141 males, 141 females, 18  
 163 litters and 43 litters where at least 50% were phenotyped) and 290 EMD with unknown  
 164 phenotype were included to form seven families. In dataset 3, pedigrees had to be adjusted  
 165 and litters in which all puppies were of the IVEU phenotype excluded (Table 3). The analysis  
 166 of dataset 1 and 2 resulted in the rejection of the environmental model. The mixed inheritance  
 167 model was superior to the general genetic model, but no difference between mixed



inheritance, major and polygene model was detected. In dataset 3, the polygene model was rejected while the major gene model was superior to the mixed inheritance model. Heritability of the EVEU trait was estimated to be 0.65 (SE  $\pm$  0.11) in a dataset of 1611 EMD including trigone (n = 196) and EVEU phenotypes (n = 115).

## Discussion

This is the first report to demonstrate a hereditary basis for EU in the dog that had been suspected for decades. A genetic involvement has been suggested with breeds of different predispositions (Hayes 1974; Holt *et al.* 2000; North *et al.* 2010) and an increased incidence of EU in the EMD has been reported previously in a clinical case study (Reichler 2012), as well as in epidemiological studies (North *et al.* 2010; Bitterli 2011). In the present study a decline in the number of EVEU affected dogs born after 2008 was observed, which coincided with the establishment of breeding restrictions based on the results of the screening program in the form of reducing or excluding EVEU affected dogs from breeding. This supports the premise of treating IVEU and EVEU as separate phenotypes and also questions the involvement of the IVEU phenotype in the development of the EVEU phenotype. Using complex segregation analysis we could show the presence of a hereditary basis of the disease. The environmental model was rejected in all three datasets and when IVEU phenotypes were excluded from the dataset the major gene model fitted best. Therefore we suggest the presence of a major gene involved with the occurrence of the EVEU phenotype in the EMD. This is further reflected in the result of a high heritability coefficient of 0.65 (SE  $\pm$  0.11) for EVEU. The power of testing may have been compromised by the fact of a nonrandom population. Most likely future breeding dogs and dogs with relatives affected by EU were overrepresented. Furthermore often only some but not all dogs of a litter were phenotyped. Even bearing the nonrandomly phenotyped population in mind it is still concerning that only one third of the animals in our study had normal ureteral terminations in the bladder trigone area and that 46% had IVEU and 19% had EVEU. In Germany and some other countries it is

194 illegal to breed animals which have a high probability of inherited disorders and welfare  
195 problems. Therefore screening examinations were introduced as mandatory for intended  
196 breeding animals and dogs with EVEU were excluded from breeding. Loosing 19% of  
197 potential breeding animals to EU does not seem like a lot, however it has to be kept in mind  
198 that in most breeds and also in the EMD only a low percentage of males are used for  
199 breeding. Together with breeding constraints already in use due to other disease conditions  
200 (hip dysplasia, progressive retina atrophy), the degree of inbreeding and thus danger to the  
201 health status of the population might increase. Discussions about outcrossing, which would be  
202 a very effective breeding strategy in such a small population having such a high incidence of  
203 EU, are ongoing. Such a breeding strategy, if performed correctly, could increase the genetic  
204 base tremendously without obviously changing outward appearance and breed characteristics.  
205 However these methods are not accepted by the FCI or the involved kennel clubs. Another  
206 option to decrease EU prevalence in this breed is the establishment of a breeding strategy  
207 using estimated breeding values. This strategy was therefore recommended to kennel clubs  
208 for future breeding.

209 The distinction between normal openings in the trigone and IVEU openings with sonography  
210 is difficult, as no normal values for the distance between ureterovesical junction and internal  
211 urethral orifice have been published. Based on measurements in freshly euthanized dogs (F.  
212 Degrandi, pers. comm.) and data from dogs evaluated by CT, where distances of 1.8 to 3.9 cm  
213 were reported (Rozear & Tidwell 2003), we considered distances greater than 1.5 to 2 cm to  
214 be normal according to the size of the dog. We accounted for the debatable identification of  
215 the IVEU phenotype in the complex segregation analysis by grouping it as either affected,  
216 unaffected or no information in the different datasets.

217 According to the results of the mixed multivariate logistic regression models the female  
218 EMD, surprisingly had a lower chance of being affected than males. Possible explanations  
219 like early loss due to resorption of embryos or neonatal death of affected females could be

220 ruled out by analyzing the ratio of female and male EMD born in the German and Swiss  
 221 populations in the last 12 years and the gender distribution of early-death puppies. Both were  
 222 similar. Bias could have arisen during diagnostic work-up if males were more strictly  
 223 classified than females. Historically, more females are reported to have the disease (Holt &  
 224 Moore 1995) although the true prevalence in males must have been underestimated (Berent *et*  
 225 *al.* 2008; Reichler *et al.* 2012). Using the prostate as a landmark in males facilitates the  
 226 classification of EVEU. In older intact male dogs benign prostatic hyperplasia has to be taken  
 227 into account, because a cranially enlarged prostate can compress the bladder neck, which may  
 228 then be mistaken for the urethra and could have biased our results. To avoid misinterpretation  
 229 during the screening process we focused therefore on the junction of the prostate and the  
 230 urethra, which concurs anatomically with the openings of the deferent ducts. Furthermore the  
 231 preliminary results of Degrandi in dog breeds not affected by EU make such a bias less likely.  
 232 Comparing intraindividually the ultrasonographic measurements of the distances between  
 233 ureterovesical junction and internal urethral orifices with the measurements post dissection,  
 234 she found neither diverging results regarding the measurement methods nor gender related  
 235 differences (F. Degrandi, pers. comm.). An X-linked mode of inheritance, which is a further  
 236 explanation for the gender distribution, could unfortunately not be ruled out due to the limited  
 237 size of our dataset and PAP requiring to estimate a larger number of parameters in this mode.  
 238 Year of birth and birth after 2008 were found to be significant covariates in the versions (ii  
 239 and iii) which exclusively assign EVEU phenotypes as affected. This reflects our observation  
 240 of a decrease of EVEU affected dogs over time, which is most likely due to the applied  
 241 breeding restrictions. The covariate age at diagnosis was found to decrease the log odds ratio  
 242 in version (ii), indicating that older dogs were less likely to have an affected phenotype in this  
 243 version. Since 2008 the screening evaluation became mandatory for all breeding dogs in  
 244 Germany and was highly encouraged in Switzerland. Supposedly a higher proportion of  
 245 young dogs but also older breeding dogs that had not shown any signs of disease were

**Kommentar [R1]:** and

**Kommentar [RF2]:** Wie vorgeschlagen

**Kommentar [R3]:** loss due to resorption of embryos or neonatal death of affected females were ruled out by analyzing the ratio of female and male EMD born in the German an Swiss populations in the last 12 years and the gender distribution of early-death puppies, which were both equal

Sorry, da war mein Vorschlag unausgereift- und ich bin auch jetzt nicht sicher ob das gut formuliert ist

**Kommentar [R4]:** Ich bin nicht ganz glücklich, Vielleicht: An X-linked mode of inheritance could explain the observed gender distribution. An attempt to proof or rule out this mode of inheritance however, was unfortunately not possible with our dataset as PAP requires to estimate a larger number of parameters in this mode and therefore a lot more phenotypes. Oder einfach die PAPA erklärung doch weglassen- wie DU willst-

**Kommentar [RF5]:** Verständlich?

**Kommentar [RF6]:** Denk ist besser so super!!

246 presented for examination. The covariate season of birth between June and August increased  
247 the risk of having an affected phenotype in version (ii) and (iii), indicating a possible non  
248 genetic influence on the affected outcome EVEU.

249 In the current study we were able to demonstrate a hereditary basis of EU in dogs by example  
250 of the EMD. The complex inheritance pattern of the disease likely involves several genes as  
251 well as a major gene in association with the clinically more relevant EVEU phenotype. This  
252 breed seems to be highly affected both clinically and phenotypically (Bitterli 2011). This may  
253 allow us to identify molecular markers that will help to understand the genetic base and  
254 pathogenesis of the disease in other dog breeds and in humans as well. Fortunately, the  
255 incidence of EVEU was observed to have decreased in the EMD shortly after breeding  
256 restrictions were implemented.

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#### 261 References

- 262 Akaike H. (1974) A new look at the statistical model identification. *IEEE Trans. Automat.*  
263 *Contr.*, 19, 716–723.
- 264 Berent A. C., Mayhew P. D., Porat-Mosenco Y. (2008) Use of cystoscopic-guided laser  
265 ablation for treatment of intramural ureteral ectopia in male dogs: four cases (2006-2007). *J.*  
266 *Am. Vet. Med. Assoc.*, 232, 1026–1034.
- 267 Bitterli F. (2011) Prävalenz und klinische Relevanz ektopischer Ureteren beim Entlebucher  
268 und Appenzeller Sennenhund. Dissertation, Vetsuisse Faculty, University of Zurich, Zurich.
- 269 Boldman, K. G., L. A. Kriese, L. D. Van Vleck, C. P. Van Tassell and S. D. Kachman (1995)  
270 A Manual for Use of MTDFREML. A Set of Programs To Obtain Estimates of Variances and

271 Covariances [DRAFT]. U.S. Department of Agriculture, Agricultural Research Service, Clay  
 272 Center.

273 Cannizzo K. L., McLoughlin M. A., Mattoon J. S., Samii V. F., Chew D. J., DiBartola S. P.  
 274 (2003) Evaluation of transurethral cystoscopy and excretory urography for diagnosis of  
 275 ectopic ureters in female dogs: 25 cases (1992-2000). *J. Am. Anim. Hosp. Assoc.*, 223, 475–  
 276 481.

277 Dean P. W., Bojrab M. J., Constantinescu G. M. (1988) Canine ectopic ureter. *Compend.*  
 278 *Contin. Educ. Vet.*, 10, 146–162.

279 Deweerd J. H., Feeney D. P. (1967) Bilateral ureteral ectopia with urinary incontinence in a  
 280 mother and daughter. *J. Urol.*, 98, 335–337.

281 Eckrich Specker C. (2006) Ektopische Ureteren beim Hund: Eine retrospektive Analyse von  
 282 30 Fällen. Dissertation, Vetsuisse Faculty, University of Zurich, Zurich.

283 Hasstedt S.J. (2002) Pedigree Analysis Package, 5.ed., Salt Lake City: Department of Human  
 284 Genetics, University of Utah, Salt Lake City.

285 Hayes H. M. Jr. (1974) Ectopic ureter in dogs: epidemiologic features. *Teratology*, 10, 129–  
 286 132.

287 Hayes H. M. Jr. (1984) Breed associations of canine ectopic ureter: a study of 217 female  
 288 cases. *J. Small Anim. Pract.*, 25, 501–504.

289 Ho L. K., Troy G. C., Waldron D. R. (2011) Clinical outcomes of surgically managed ectopic  
 290 ureters in 33 dogs. *J. Am. Anim. Hosp. Assoc.*, 47, 196–202.

291 Holt P. E., Gibbs C., Pearson H. (1982) Canine ectopic ureter – a review of twenty-nine cases.  
 292 *J. Small Anim. Pract.*, 23, 195–208.

293 Holt P. E. (1990) Urinary incontinence in dogs and cats. *Vet. Rec.*, 127, 347–350.

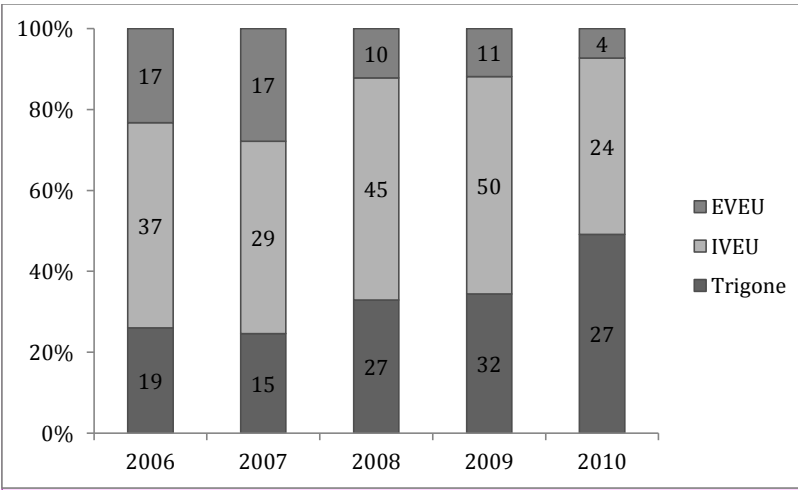
294 Holt P. E., Moore A. H. (1995) Canine ureteral ectopia: an analysis of 175 cases and  
 295 comparison of surgical treatments. *Vet. Rec.*, 136, 345.

296 Holt P. E., Thrusfield M. V., Hotston Moore A. (2000) Breed predisposition to ureteral  
 297 ectopia in bitches in the UK. *Vet. Rec.*, 146, 561.  
 298 Johnston G. R., Osbourne C. A., Wilson J. W., Yano B. L. (1977) Familial Ureteral Ectopia in  
 299 the Dog. *J. Am. Anim. Hosp. Assoc.*, 13, 168–170.  
 300 Lamb C. R., Gregory S. P. (1998) Ultrasonographic findings in 14 dogs with ectopic ureter.  
 301 *Vet. Radiol. Ultrasound*, 39, 218–223.  
 302 Musselman B. C., Barry J. J. (1973) Varying degrees of ureteral ectopia and duplication in 5  
 303 siblings. *J. Uro.*, 110, 476–477.  
 304 North C., Kruger J. M., Venta P. J., Miller J. M., Rosenstein D. S., Randall E. K., White B.,  
 305 Fitzgerald S. D. (2010) Congenital ureteral ectopia in continent and incontinent-related  
 306 Entlebucher mountain dogs: 13 cases (2006–2009). *J. Vet. Intern. Med.*, 24, 1055–1062.  
 307 Osborne C. A., Johnston G. R., Kruger J. M. (1995) Ectopic ureters and ureteroceles. In:  
 308 Osborne C. A., Finco D. R. (eds.) *Canine and feline nephrology and urology*. 1st ed.  
 309 Philadelphia: Williams & Wilkins, 608–622.  
 310 Reichler I. M., Eckrich Specker C., Hubler M., Boos A., Haessig M., Arnold S. (2012)  
 311 Ectopic Ureters in Dogs: Clinical Features, Surgical Techniques and Outcome. *Vet. Surg.*, 41,  
 312 515–522.  
 313 Rozear L., Tidwell A. S. (2003) Evaluation of the ureter and ureterovesicular junction using  
 314 helical computed tomographic excretory urography in healthy dogs. *Vet. Radiol. Ultrasound*,  
 315 44, 155–164.  
 316 Smith C. W., Stowater J. L., Kneller S. K. (1981) Ectopic ureter in the dog: a review of cases.  
 317 *J. Am. Anim. Hosp. Assoc.*, 17, 245–248.  
 318 Snow G. L., Wijsman E. M. (1998) Pedigree analysis package (PAP) vs. MORGAN: model  
 319 selection and hypothesis testing on a large pedigree. *Genet. Epidemiol.*, 15, 355–369.  
 320 Staub K. (2011) Untersuchungen zur Rute beim Entlebucher Sennenhund. Dissertation,  
 321 Vetsuisse Faculty, University of Zurich, Zurich.

Stone E. A., Mason L. K. (1990) Surgery of ectopic ureters: types, method of correction, and postoperative results. *J. Am. Anim. Hosp. Assoc.*, 26, 81–88.

Figures

Figure 1 **Decline of affected EMD after establishment of breeding restrictions.**



**Kommentar [RF7]:** Nimms noch raus, keine Sorge!

Phenotype distribution as bilateral terminations in the vesical trigone, IVEU (at least one ureteral opening is intravesically ectopic) or EVEU (at least one ureteral opening is extravesically ectopic) in EMD born two years before and after breeding restrictions were applied in 2008. The absolute number of phenotyped dogs is given within bars.

Tables

Table 1 Year of birth and gender distribution of 565 Entlebucher Mountain Dogs examined for ectopic ureters and phenotyped as bilateral terminations in the vesical trigone, IVEU (at least one ureteral opening is intravesically ectopic), EVEU (at least one ureteral opening is extravesically ectopic) or unknown.

Year of birth	Total Male/Female	Phenotype trigone Male/Female	Phenotype IVEU Male/Female	Phenotype EVEU Male/Female	Phenotype unknown Male/Female
Before 2002	40 27/13	12 7/5	8 2/6	17 16/1	3 2/1
2002	24 14/10	4 1/3	11 6/5	8 6/2	1 1/0

2003	28 12/16	9 1/8	12 7/5	7 4/3	0
2004	47 19/28	16 6/10	22 7/15	6 4/2	3 2/1
2005	48 26/22	18 8/10	19 10/9	11 8/3	0
2006	73 36/37	19 3/16	37 19/18	17 14/3	0
2007	63 36/27	15 5/10	29 18/11	17 13/4	2 0/2
2008	86 37/49	27 7/20	45 21/24	10 7/3	4 2/2
2009	93 45/48	32 9/23	50 29/21	11 7/4	0
2010	55 19/36	27 7/20	24 8/16	4 4/0	0
2011	8 6/2	2 0/2	5 5/0	1 1/0	0
Total	565 277/288	181 54/127	262 132/130	109 84/25	13 7/6

336

337 Table 2 Results of the mixed multivariate logistic regression models for EU phenotypes in  
338 EMD: Implementation of the phenotypes trigone (bilateral terminations in the vesical  
339 trigone), IVEU (at least one ureteral opening is intravesically ectopic) and EVEU (at least one  
340 ureteral opening is extravesically ectopic) with binary encoding in three versions: (i) trigone  
341 as unaffected, IVEU together with EVEU as affected; (ii) trigone together with IVEU as  
342 unaffected, EVEU as affected; and (iii) trigone as unaffected and EVEU as affected while  
343 specifying IVEU as no information.

Covariates	Version (i)		Version (ii)		Version (iii)	
	Log odds ratio	P -value	Log odds ratio	P -value	Log odds ratio	P -value
Gender (female)	- 1.673	< 0.001	- 1.776	< 0.001	- 2.996	< 0.001
Sex ratio	- 0.678	0.213	- 0.813	0.200	- 1.189	0.173
Birth after 2008	- 0.363	0.188	- 0.232	0.644	- 1.227	0.037
Year of birth			- 0.45	0.002		
Season 2			0.585	0.219	0.709	0.263
Season 3			1.141	0.015	1.759	0.006
Season 4			0.481	0.345	0.776	0.262
Age at diagnosis			- 0.384	0.014	- 0.002	0.981

344



Table 3 Segregation analysis of trichotomous (T) and dichotomous (D) datasets from different phenotype groupings for ectopic ureters in Entlebucher Mountain Dogs. Prevalences are given according to the number of phenotyped animals for trigone, IVEU (Intravesical ectopic ureter) and EVEU (Extravesical ectopic ureter) phenotypes. Chi square values ( $\chi^2$ ) are equal to the difference of the environmental and mixed inheritance model tested with the general genetic model or of the major gene and polygene model tested with the mixed inheritance model, respectively. Degrees of freedom (df) are equal to the difference in estimated parameters between compared models.

Dataset	Phenotype	Prevalence	Animals	Model comparison		$\chi^2$	df	P-value
1	Trigone	0.33	Total 572 Phenotyped 282	General genetic	- Environmental	11.947	4	0.018
					- Mixed inheritance	2.283	3	0.516
	IVEU and EVEU	0.67			- Major gene	0.199	1	0.656
					- Polygene	1.897	3	0.594
2	Trigone and IVEU	0.80	Total 572 Phenotyped 282	General genetic	- Environmental	21.390	4	< 0.001
					- Mixed inheritance	1.755	3	0.625
	EVEU	0.20			- Major gene	2.911	1	0.088
					- Polygene	5.534	3	0.130
3	Trigone	0.80	Total 451 Phenotyped 137	General genetic	- Environmental	26.012	4	< 0.001
					- Mixed inheritance	4.530	3	0.210
	EVEU	0.20			- Major gene	0.452	1	0.501
					- Polygene	20.517	3	< 0.001